



1600

RAW SEQUENCE LISTING

DATE: 07/14/2003

PATENT APPLICATION: US/09/913,351A

TIME: 11:55:44

Input Set : A:\618JPB75-corrected.txt

Output Set: N:\CRF4\07142003\I913351A.raw

3 <110> APPLICANT: DELGADO, AURORA BRIEVA
 4 VILLARRUBIA, VINCENTE GARCIA
 5 GOMEZ-PAMO, ANTONIO GUERRERO
 6 RANIERI, JUAN PABLO PIVEL
 7 GALLEGO, GUILLERMO GIMENEZ
 8 TUDURI, JOSE ANTONIO MATJI
 10 <120> TITLE OF INVENTION: PHARMACOLOGICALLY ACTIVE POLYPEPTIDE GLYCOCONJUGATES
 12 <130> FILE REFERENCE: 618999-1/JP/B-4275
 14 <140> CURRENT APPLICATION NUMBER: 09/913,351A
 15 <141> CURRENT FILING DATE: 1999-10-21
 17 <150> PRIOR APPLICATION NUMBER: PCT/ES99/00338
 18 <151> PRIOR FILING DATE: 1999-10-21
 20 <150> PRIOR APPLICATION NUMBER: ES P9900408
 21 <151> PRIOR FILING DATE: 1999-02-26
 23 <160> NUMBER OF SEQ ID NOS: 5
 25 <170> SOFTWARE: PatentIn Ver. 2.1

ERRORED SEQUENCES

246 <210> SEQ ID NO: 5
 247 <211> LENGTH: 65
 248 <212> TYPE: PRT
 249 <213> ORGANISM: Ricinus communis
 251 <400> SEQUENCE: 5
 252 Gln Glu Arg Ser Leu Arg Gly Cys Cys Asp His Leu Lys Gln Met Gln
 253 1 5 10 15
 255 Ser Gln Cys Arg Cys Glu Gly Leu Arg Gln Ala Ile Glu Gln Gln Gln
 256 20 25 30
 258 Ser Gln Gly Gln Leu Gln Gly Gln Asp Val Phe Glu Ala Phe Arg Thr
 259 35 40 45
 261 Ala Ala Asn Leu Pro Ser Met Cys Gly Val Ser Pro Thr Glu Cys Arg
 262 50 55 60
 264 Phe
 265 55

Does Not Comply
Corrected Diskette Needed

E--> 268 5

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/913,351A

DATE: 07/14/2003

TIME: 11:55:45

Input Set : A:\618JPB75-corrected.txt

Output Set: N:\CRF4\07142003\I913351A.raw

L:146 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
M:341 Repeated in SeqNo=1
L:268 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:5

STATISTICS SUMMARY

PATENT APPLICATION: US/09/913,351A

DATE: 07/14/2003

TIME: 11:55:45

Input Set : A:\618JPB75-corrected.txt

Output Set: N:\CRF4\07142003\I913351A.raw

Application Serial Number: US/09/913,351A

Alpha or Numeric or Xml: Numeric

Application Class:

Application File Date: 10-21-1999

Art Unit: 1600

Software Application: PatentIN2.1

Total Number of Sequences: 5

Total Nucleotides: 0

Total Amino Acids: 434

Number of Errors: 1

Number of Warnings: 15

Number of Corrections: 0

MESSAGE SUMMARY

332 E: 1 ((32) Invalid/Missing Amino Acid Numbering)

341 W: 15 ((46) "n" or "Xaa" used)